

1  
GAATTCCGAT TTAGCCTCAT ACTGCTTCTC ACATTACATT GGGATGCGCT  
51  
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101  
GAAGCAAACC CTTAAGAACG TCACAAACTA CATTACTGAT ATCATCTGCA  
151  
AGCGTGCAGA TCTTGGTTAC AACTATGGGG TTATCCTTAT ACCAGAAGGC  
201  
CTGATTGATT TCATCCCAGA GGTTCAAAAAA CTCATCGCAG AATTGAATGA  
251  
AATTTGGCA CATGATGTGG TTGATGAGGC AGGGGCCTGG AAAAGCAAGC  
301  
TTCAGCCTGA ATCAAAGGAG CTGTTGAGT TTTGCCCAA AACTATTAG  
351  
GAGCAACTTA TGCTTGAAAG GGGCCCCAT GGCAATGTTG AGGTTGCAAA  
401  
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451  
AGAGAAAAGC AGAGGGGAGA TACTCTGCAC ATTCAGAGG GCAAGCTCAT  
501  
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551  
CTATTGCTAT GCATTAGGCT ATGGTGCTGG TGCCCTTCTC CAAAGTGGGA  
601  
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651  
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701  
GCATGGCAAG TTCAAGCCAG TGATCAAGAA GGCTATGGTG GAACTTGATG  
751  
CTGCACCTT CAAGAAATAT GCATCAATGC GGGATGAGTG GGCCACCAAG  
801  
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851  
TGACTCGAAC CACACTTGA TGCTGGAAC CGGTGCTGAG TTATAGAGAT  
901  
GCGTCCTTG CTTATTTTG TTTCTTACAG TTTGGGAGT GGAGACTGGA  
951  
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1001  
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1051  
AGAACTGGTT TTAGCATTAA TTGTATGATT TACGCACCAA CTGACTTGTC  
1101  
TTGTAACCCCT GATTCTGTTCACTGGTTGC /ATCTCGTGA GAATGAACAA  
1151  
GTTGATATGA GGCTAAATCG GAATTC

Figure 1.

1  
ATGGCGGCAG CGAGCGGACC ATCACCTGGG ACTGGGAGGT TGGCGTCGGT  
51  
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101  
CCGTCCTCTG CTCCCAATT CTCCTCGTCG ATGGACCTCC CAGCTCAGCC  
151  
ACGGGAAACC CGGATGAGAT CGCGAAGCTG TTCCCTAACT TGTTTGGCA  
201  
GCCGTCGGCG ACATTGGTGC CGGCCAAAGA GGC GGTTGGAG GGGAAAGGCGC  
251  
TGAAGGTCGG GGTGGTGCTC TCTGGTGAC AAGCACCCGG TGGGCACAAT  
301  
GTGATCTGCG GTATCTCGA TTTCTTGAG AACACCGCAA AGGGAAAGCAC  
351  
AATGTATGGA TTCAAAGGAG GCCCAGCAGG GGTGATGAAG TGCAAGTACG  
401  
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451  
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501  
GCAAGCCGAA GATAAGCCA ACAAAACTTGA GTTGGACGGA CTTGTTGTTA  
551  
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601  
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651  
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701  
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751  
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801  
TCACATTACA TTGGGATGCG CTTGCAAAC ACACCCCAAT GCTGCACTCA  
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901  
TACATTACTG ATATCATCTG CGAGCGTGCA GATCTGGTT ACAACTATGG  
951  
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1001  
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1051  
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1101  
GTTTTGCCA AAAACTATTG AGGAGCAACT TATGCTTGAA AGGGGCC  
1151  
ATGGCAATGT TCAGGTTGCA AAAATTGAAA CCGAGAAAAT GCTTATTAGC  
1201  
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Figure 2

1251  
ACATTCAGA GGGCAAGCTC ATTTCTTGG GTACGAAGGA AGATGTGGCC  
1301  
TTCCCTACCAA TTTTGATTCT AACTATTGCT ATGCATTAGG CTATGGGCT  
1351  
GGTGCCCTTC TCCAAAGTGG GAAGACAGGA CTTATTCAT CGGTTGGCAA  
1401  
CCTTGCGGCT CCAGTAGAAG AATGGACTGT TGGTGGAACAGCATTGACAT  
1451  
CACTGATGGA TGTGGAGAGG AGGCATGGCA AGTTCAAGCC AGTGATCGAG  
  
1501  
AAGGCTATGG TGGAACCTGA TGCTGCACCT TTCAAGAAAT ATGCATCAAT  
1551  
GGGGGATGAG TGGGCCACCA AGAACAGATA CATCAGCCCT GGCCCCATCC  
1601  
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Figure 2 cont.

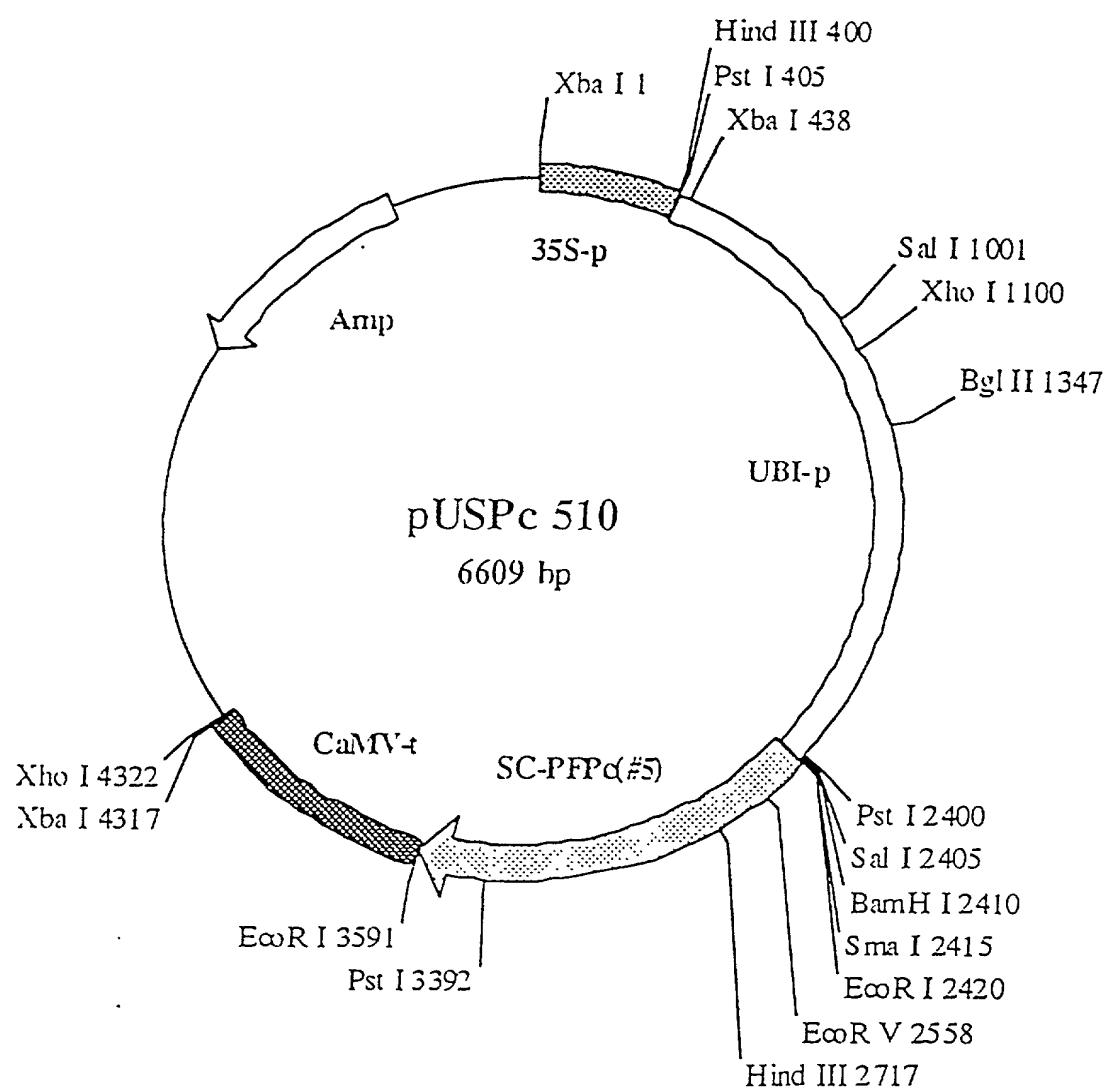


Figure 3

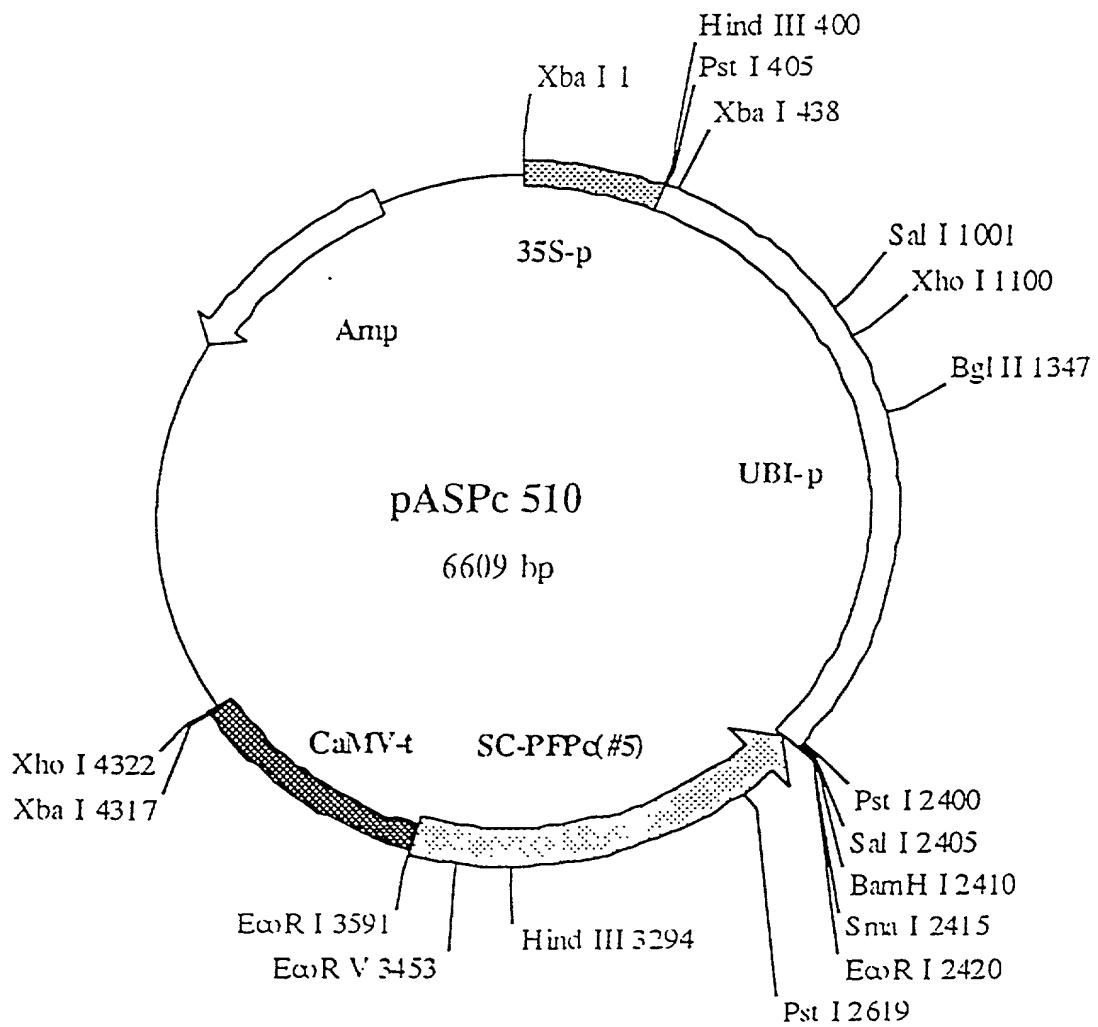


Figure 4

1 2 3 4 5 6 7 8 9 10 11 12



Figure 5 .

1 2 3 4 5

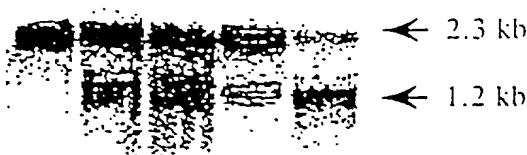


Figure 6

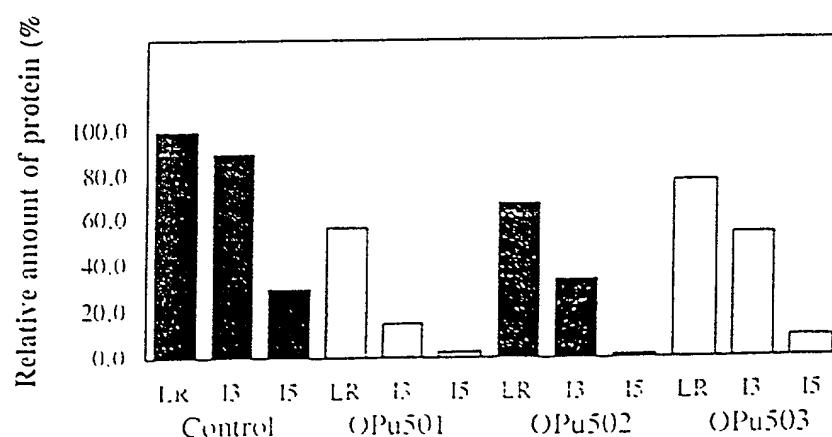


Figure 7